

# **pTNMAX (general vector)**

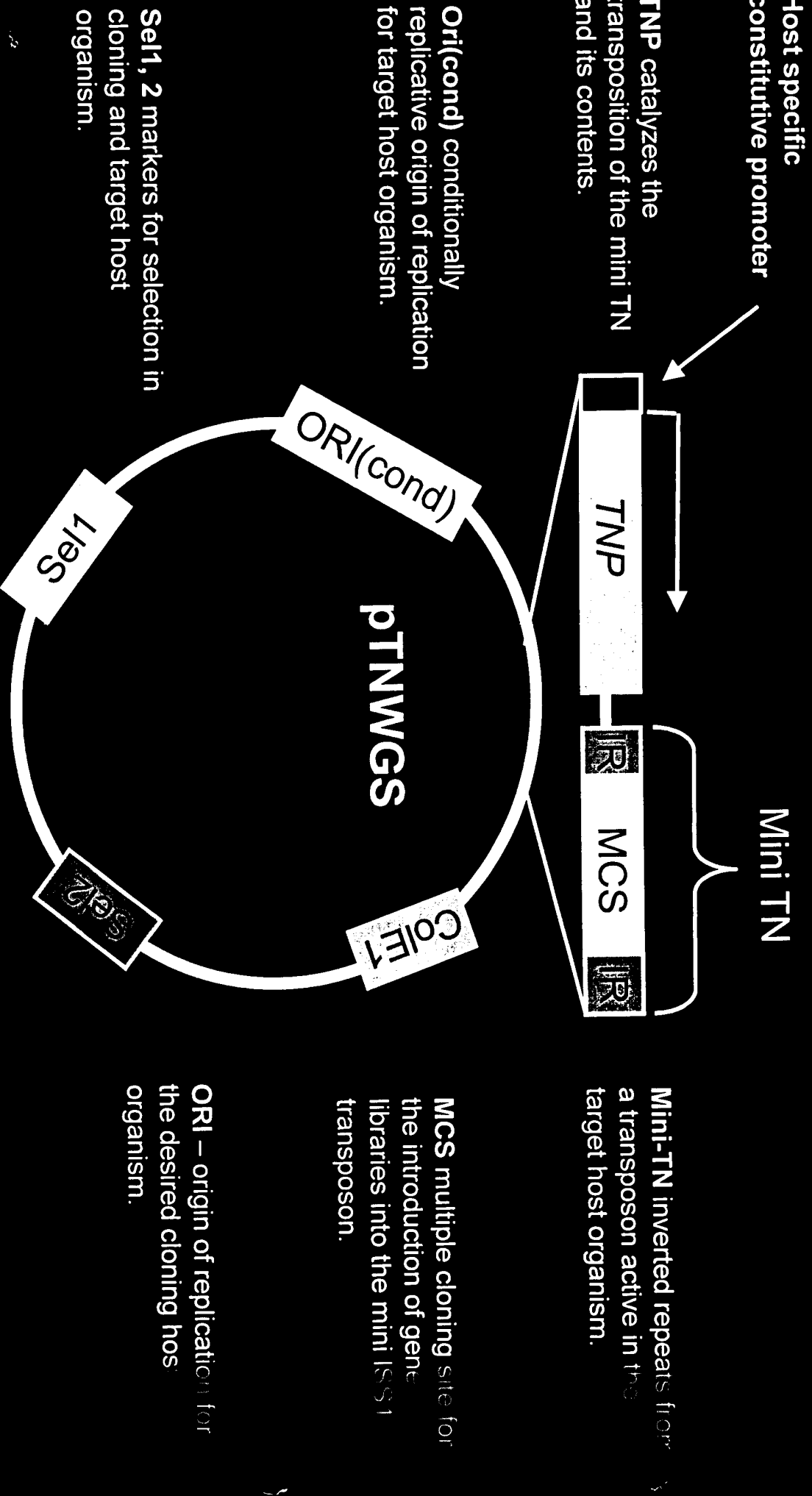
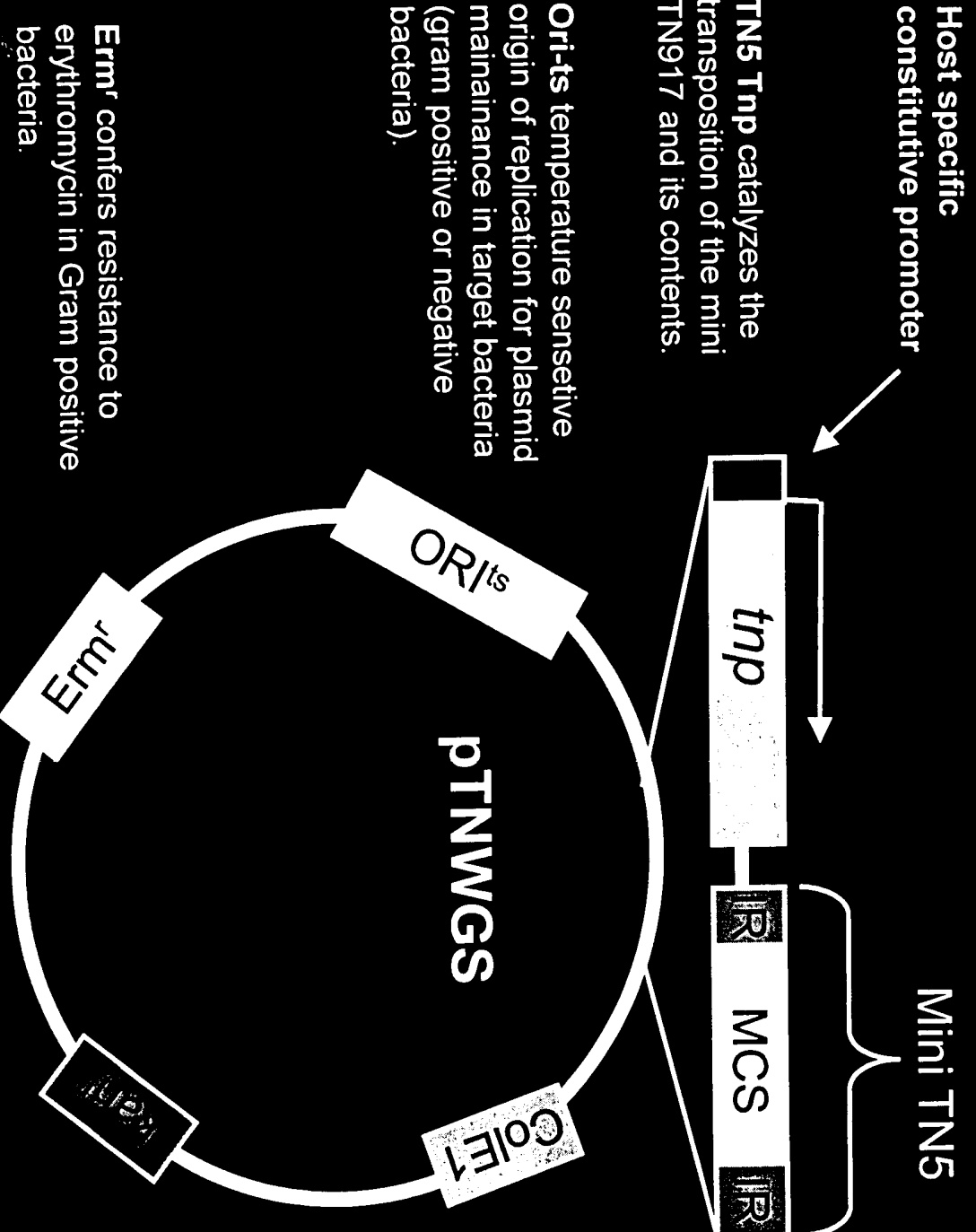


Figure 1 A

# pWGS:5



Mini TN5 TN5 inverted repeats flanking a multiple cloning site into which gene libraries can be cloned.

ColE1 origin of replication for plasmid maintenance in *E. coli*.

Kan<sup>r</sup> confers resistance to kanamycin to *E. coli*

Erm<sup>r</sup> confers resistance to erythromycin in Gram positive bacteria.

Ori<sup>ts</sup> temperature sensitive origin of replication for plasmid maintenance in target bacteria (gram positive or negative bacteria).

Figure 1B

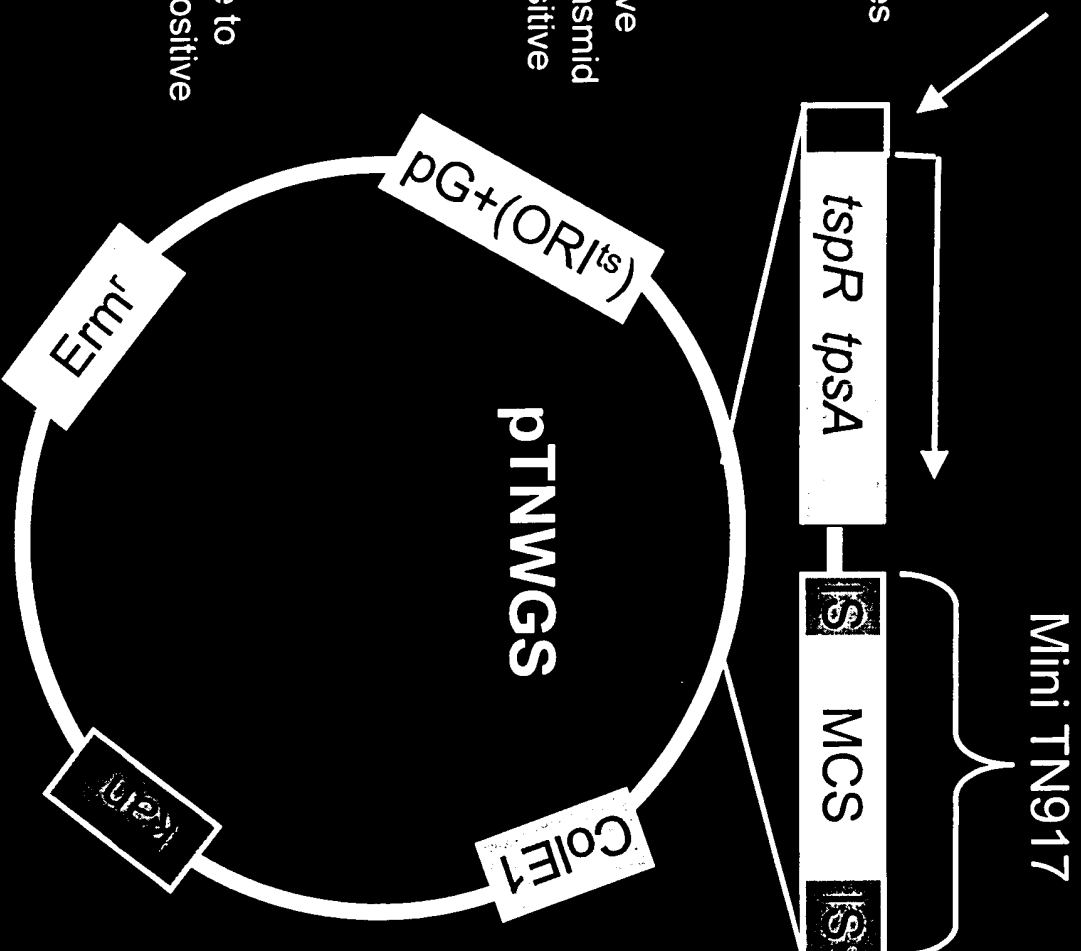
# pWGS:917

**Host specific promoter** – *nisA* promoter for lactic acid bacteria.

**917 TspR TspA** catalyzes a transposition of the mini 917 and its contents (transposase/resolvase)

**PG+** temperature sensitive origin of replication for plasmid maintenance in Gram positive bacteria.

**Erm<sup>r</sup>** confers resistance to erythromycin in Gram positive bacteria.



**MCS** multiple cloning site for the introduction of gene libraries into the mini 917 transposon.

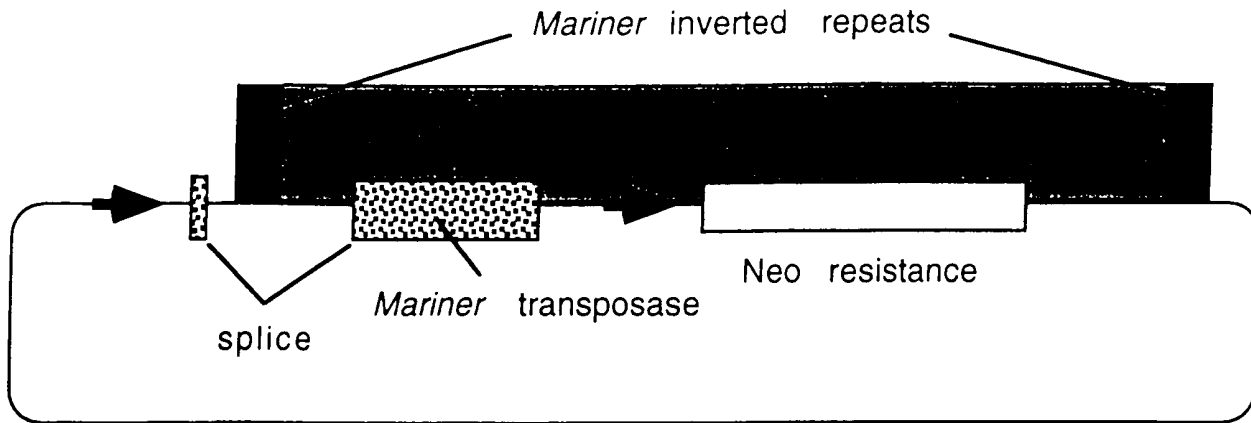
**ColE1** origin of replication for plasmid maintenance in *E. coli*.

**Kan<sup>r</sup>** confers resistance to kanamycin to *E. coli*.

Figure 1C

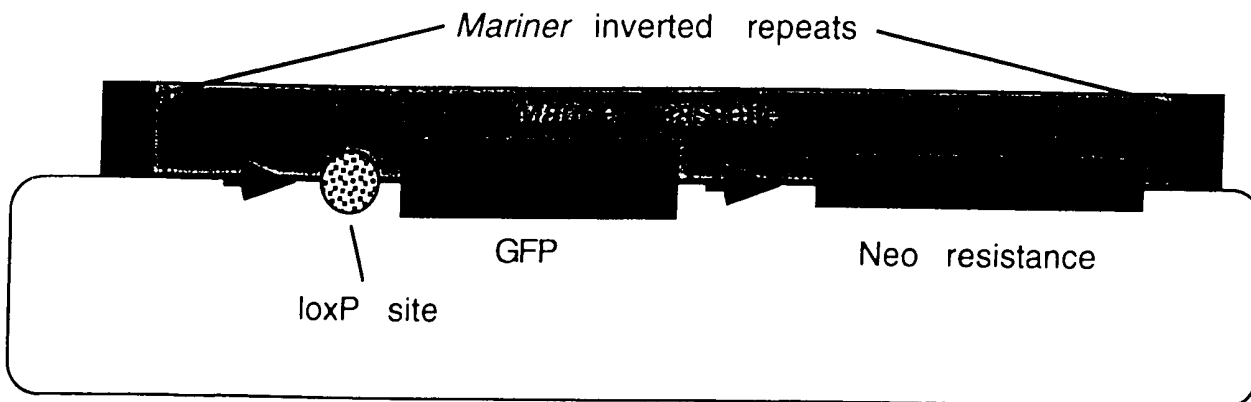
A

# Efficient integration into mammalian cells using evolved *Mariner* transposons



B

# *Mariner* transposon for inserting loxP sites at loci with desirable expression properties



# Methodology for Isolating Hosts with improved Phenotypes by Whole Genome Shuffling (WGS)

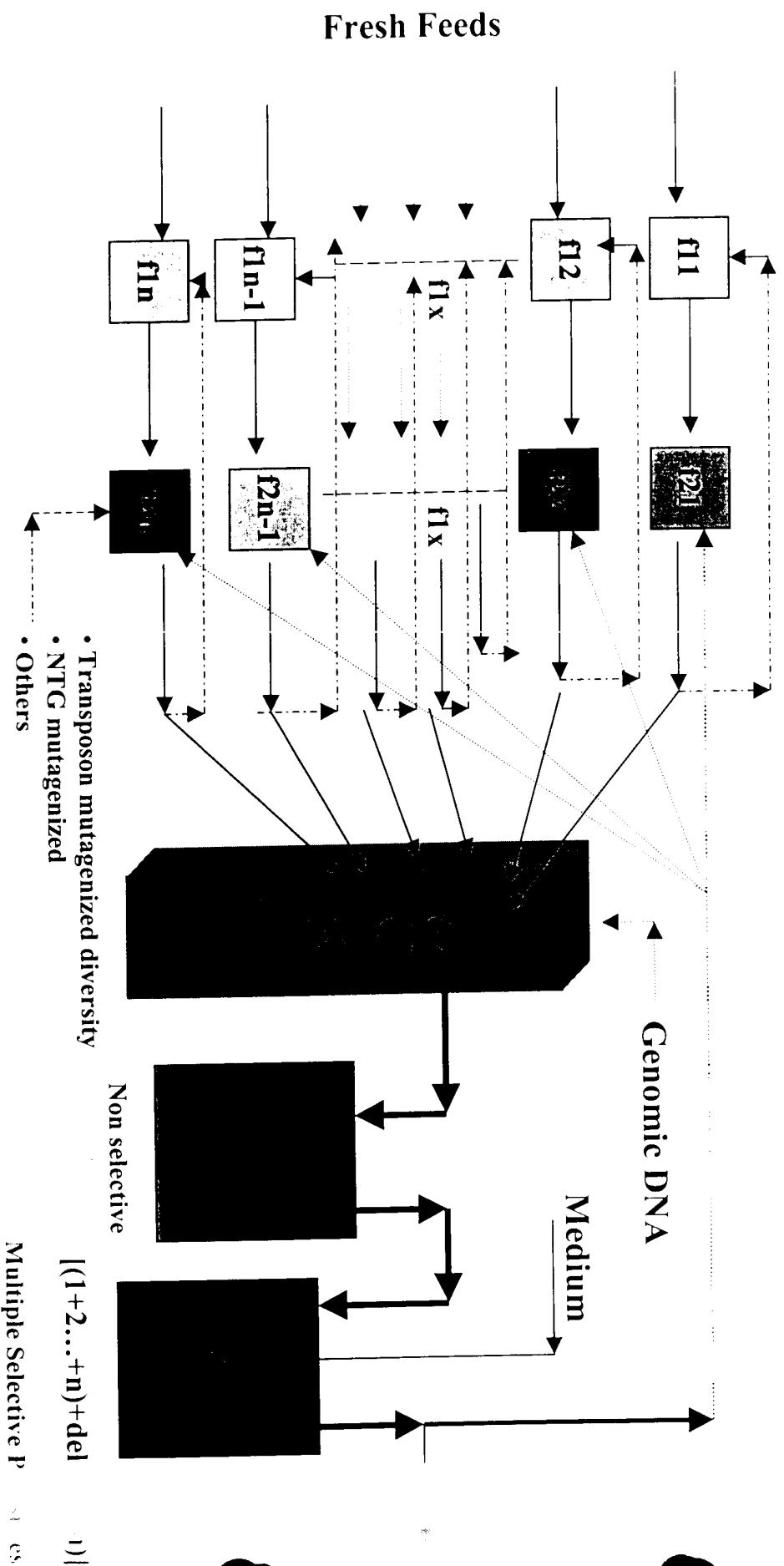


Figure 3

# Shuffling of Genomes *In Vitro*: Formation of transposomes

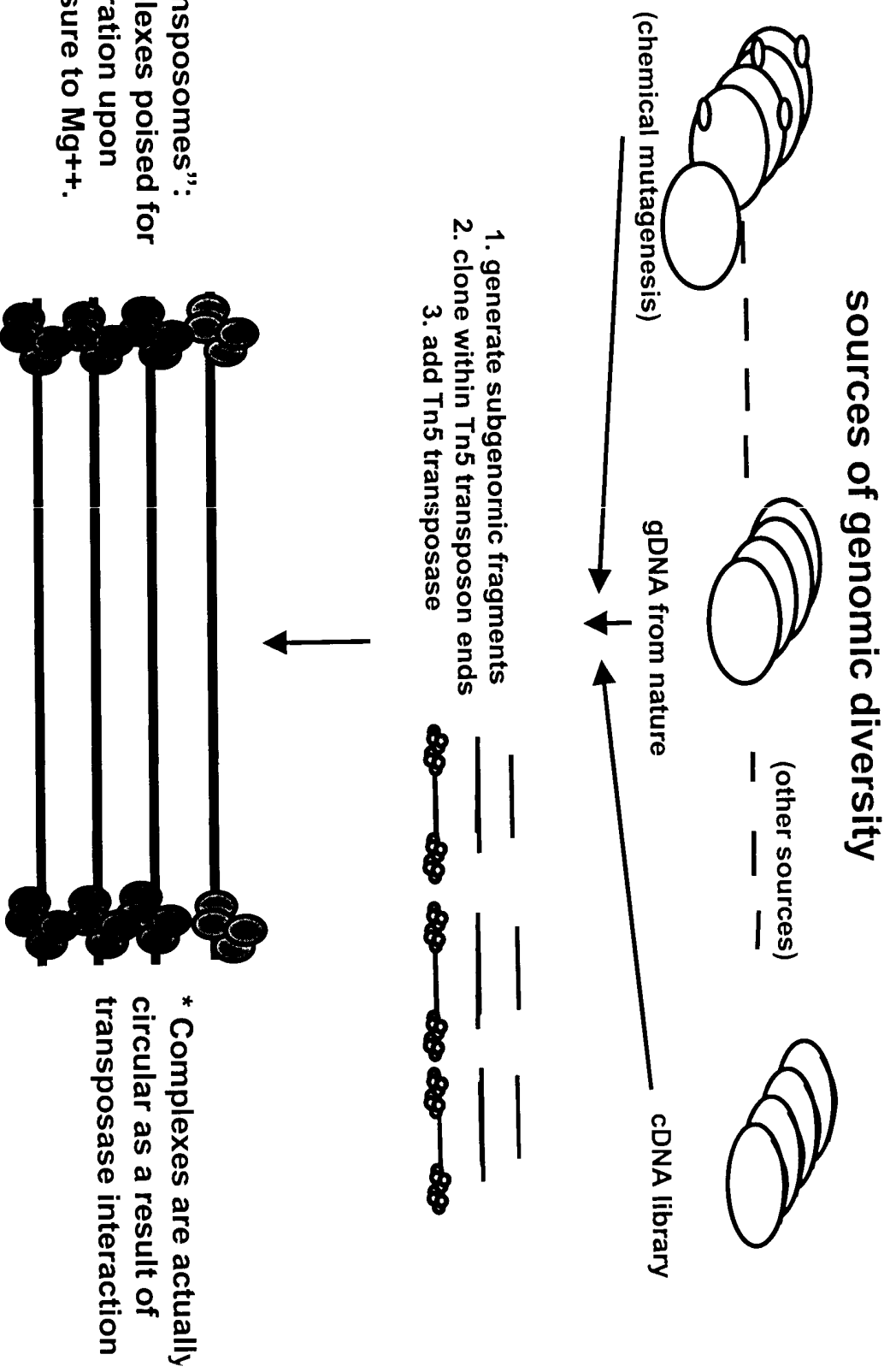


Figure 4A

# Shuffling of Genomes *In Vitro*: Breeding multiple donor genomes with a single acceptor genome

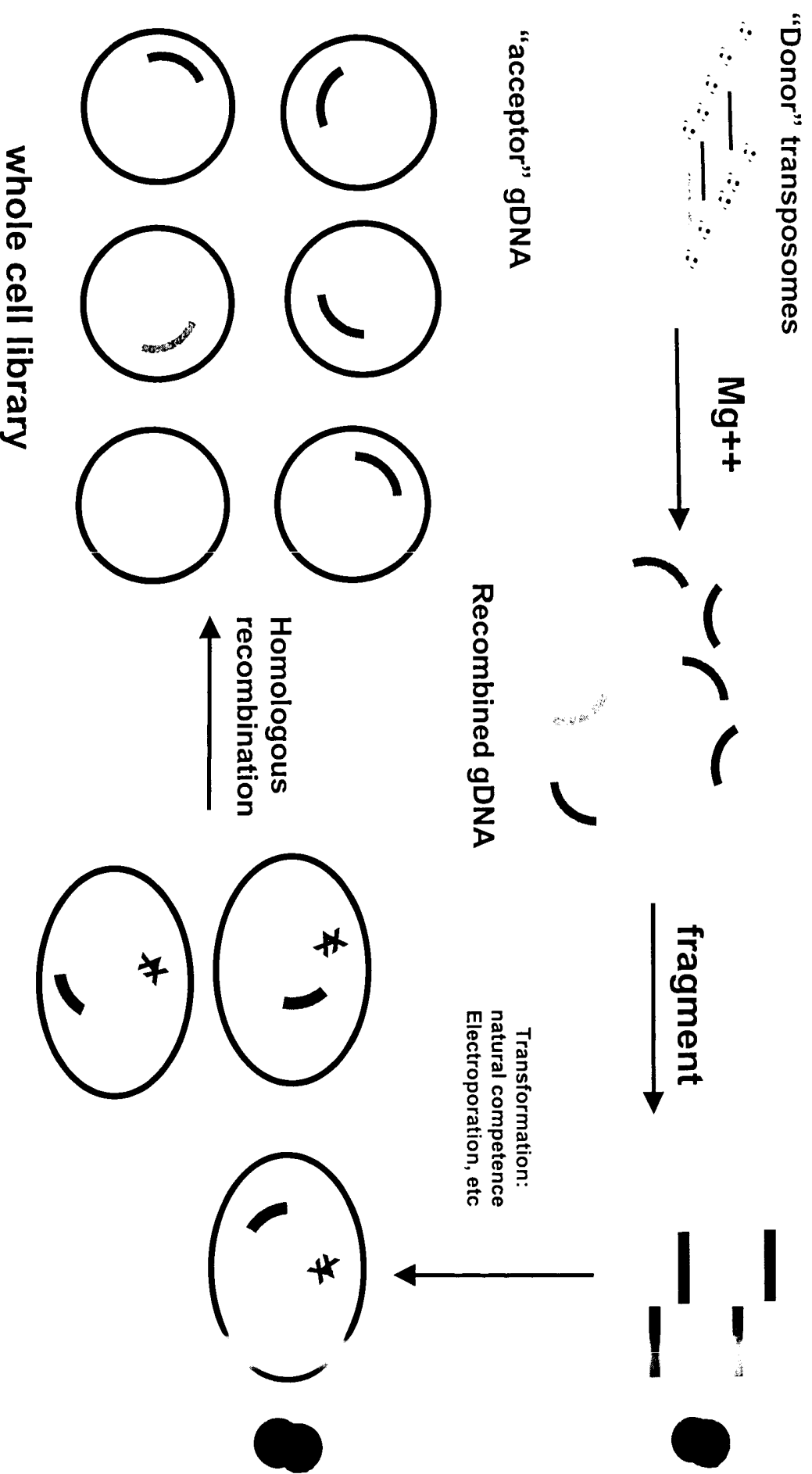


Figure 4B

# Shuffling of Genomes *In Vitro*: Breeding multiple donor genomes with multiple acceptor genomes

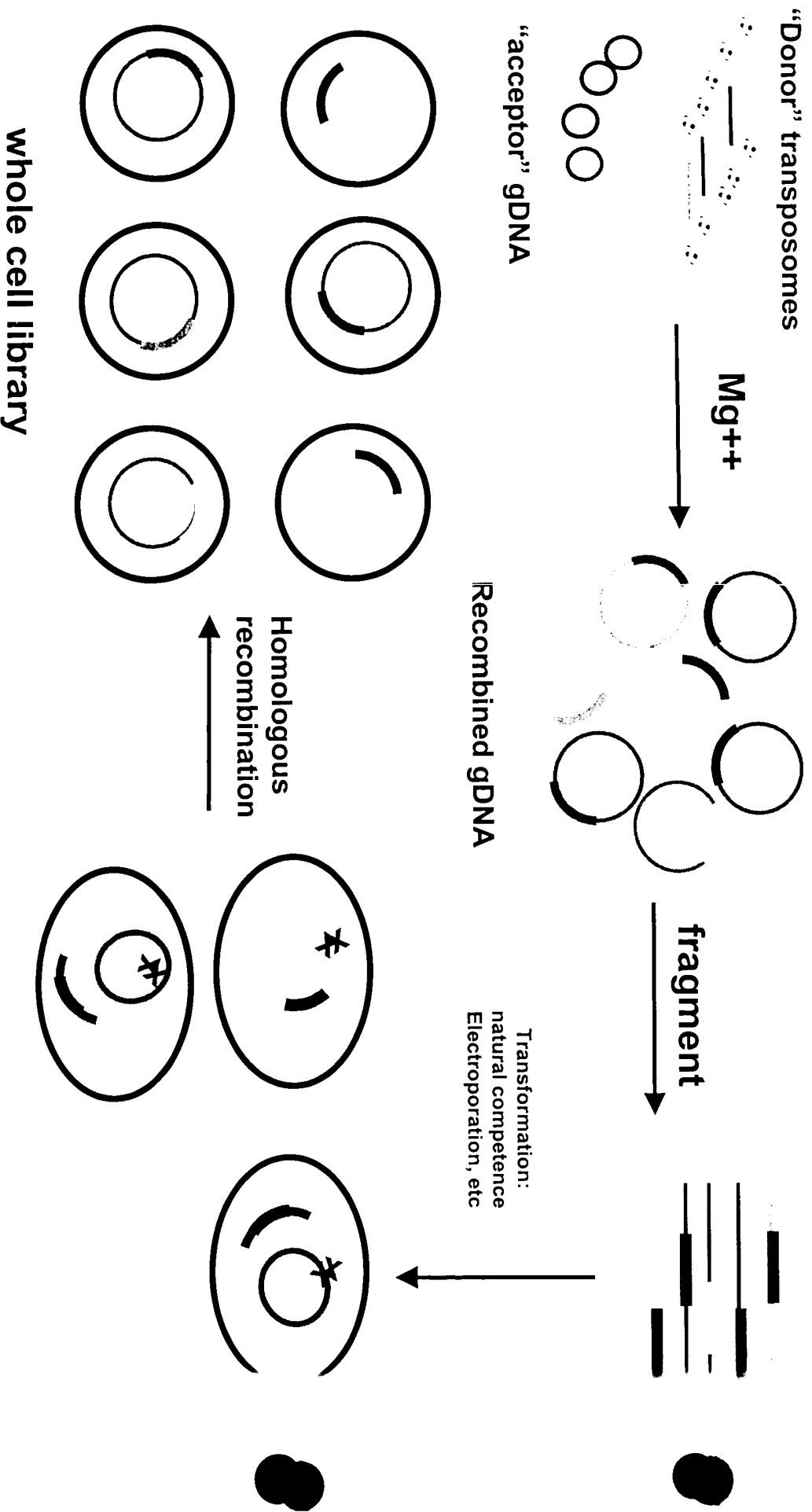


Figure 4C



# Shuffling of Genomes *In Vitro*:

Split pool recursive *in vitro* recombination of multiple genomes

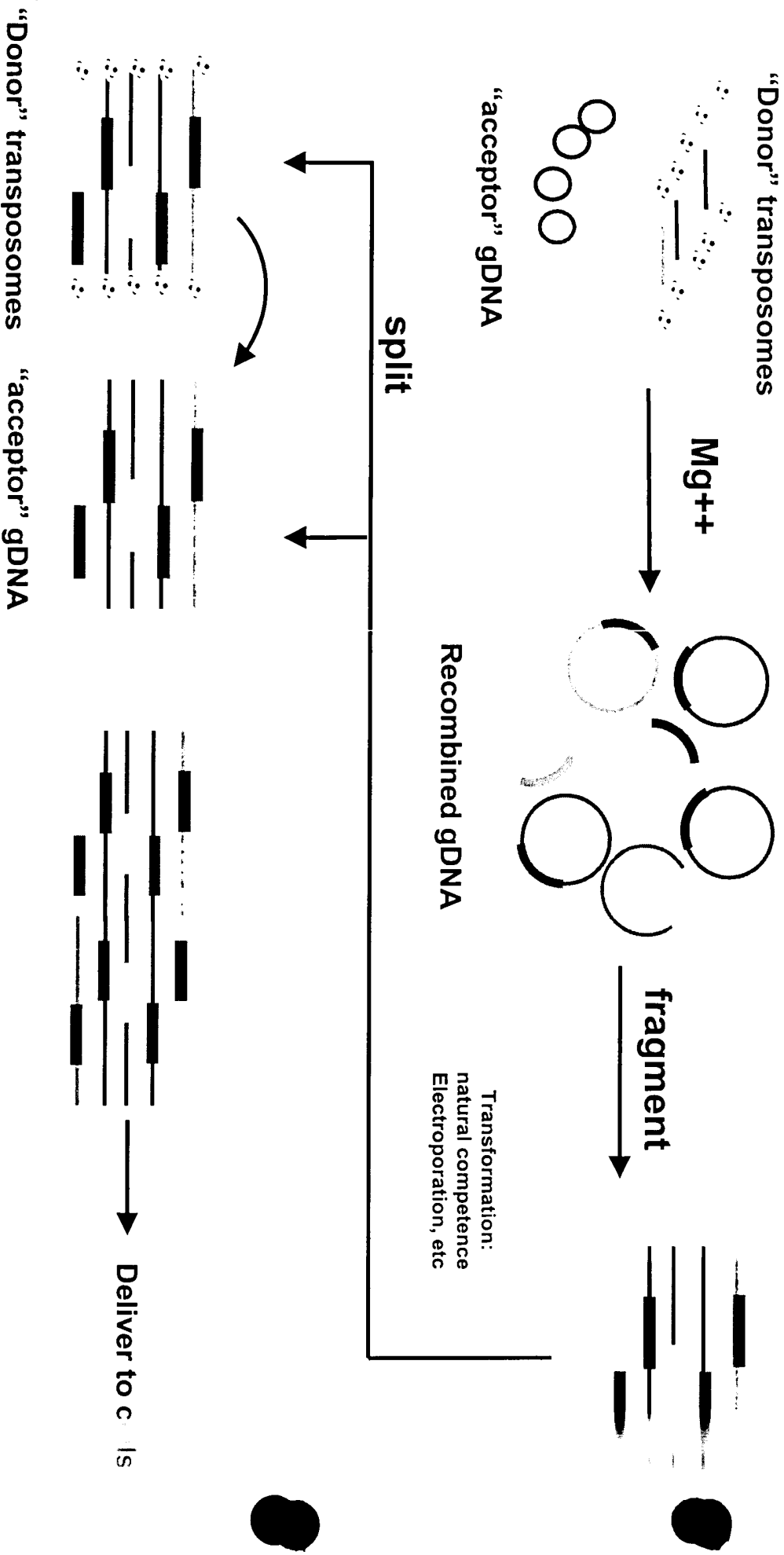


Figure 4D